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## Generic DB Entry Retrieval

ID HSM017132 standard; RNA; EST; 507 BP.  
 XX  
 AC AL121363;  
 XX  
 SV AL121363.1  
 XX  
 DT 29-SEP-1999 (Rel. 61, Created)  
 DT 20-FEB-2000 (Rel. 62, Last updated, Version 2)  
 XX  
 DE Homo sapiens mRNA; EST DKFZp762C027\_r1 (from clone DKFZp762C027)  
 XX  
 KW EST; expressed sequence tag.  
 XX  
 OS Homo sapiens (human)  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia  
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 XX  
 SN [1]  
 FP 1-507  
 FA Ottenwaelder B., Chermajer B., Mewes H.W., Gassenhuber J., Wiemann S.;  
 FT ;  
 FL Submitted (18-FEB-2000) to the EMBL/GenBank/DDBJ databases.  
 FL MIPs, Am Klopferspitz 18a, D-32152 Martinsried, GERMANY  
 XX  
 RP RZPD; DKFZp762C027; DKFZp762C027.  
 DF UNILIB; 2390; 2395.  
 XX  
 CC This is the 5' sequence of the clone insert  
 CC Clone from S. Wiemann, Molecular Genome Analysis, German Cancer  
 CC Research Center (DKFZ); Email: s.wiemann@dkfz-heidelberg.de;  
 CC sequenced by MediGenomix (Martinsried, Germany) within the cDNA  
 CC sequencing consortium of the German Genome Project.  
 CC 5' sequence also available.  
 CC This clone (DKFZp762C027) is available at the EBC in Berlin.  
 CC Please contact the RZPD: Ressourcenzentrum, Hübnerweg 6,  
 CC 14059 Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de  
 XX  
 FH Key Location/Qualifiers  
 FH  
 FT  
 FT  
 FT

FT /dev\_stage="adult"  
FT /tissue\_type="melanoma (MeWo cell line)"  
XX

Sequence 507 BP; 117 A; 152 C; 151 G; 87 T; 0 other;  
SQ  
ggctggcctc tggaaaaaagc atataactat gtaaagcaga agcggagcat cacgggcccc 6  
aacgggggct ttatgagga cgtgtatgag tatgaaggca tcttggatgc aagcaaaacag 12  
cggcacaaca agctgtggcg tcagcagaca gacagcagcc tcagcagcc tgtggatgac 18  
cctgcaggac ctggcgactt cttgcagag agcccagatg gcaccccgga aagccagctg 24  
cctctcttgg atgatgcgc ccagcccggc ttagggcccc ccttccctg ctgtttccgg 30  
cgactctcag acccccttct gccttccctt gaggatgaaa ctggcagctt ggtccacctg 36  
gaggatccgg agagggaggg tctgttggag gaagctgctc cactgcaga ggtgcacagg 42  
ccggccagac agccccagca aggttccgga ctctgtgaga aggatgtgaa gaagaaaacta 48  
gagtttggga gtcccaaaagg tgggagc 50

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